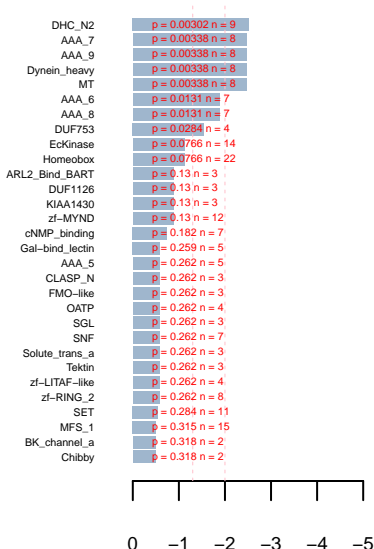


Bulk.Fem-Mal_difneg domain



log(p)

n = 1720 genes | p adjust BH TRUE